

WO 00/20588

PCT/US99/22908

SEQUENCE LISTING

<110> INCYTE PHARMACEUTICALS, INC.

TANG, Y. Tom

CORLEY, Neil C.

GUEGLER, Karl J.

LU, Aina Dyung M.

<120> BONE MARROW-DERIVED SERUM PROTEINS

<130> PF-0609 PCT

<140> To Be Assigned

<141> Herewith

<150> 09/165,621; unassigned

<151> 1998-10-02; 1999-10-02

<160> 5

<170> PERL Program

<210> 1

<211> 234

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 135698CD1

<400> 1

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Pro	Asp	Thr	Thr	Gly	Glu	Ile	Val	Leu	Thr	Gln	Ser	Pro	Ala	Thr	20	25	30	
Leu	Ser	Leu	Ser	Pro	Gly	Glu	Arg	Ala	Thr	Leu	Ser	Cys	Arg	Ala	35	40	45	
Ser	Gln	Ser	Val	Ser	Ser	Tyr	Leu	Ala	Trp	Tyr	Gln	Gln	Lys	Pro	50	55	60	
Gly	Gln	Ala	Pro	Arg	Leu	Leu	Ile	Tyr	Asp	Ala	Ser	Asn	Arg	Ala	65	70	75	
Thr	Gly	Ile	Pro	Pro	Arg	Phe	Ser	Gly	Ser	Gly	Ser	Gly	Thr	Asp	80	85	90	
Phe	Thr	Leu	Thr	Ile	Ser	Arg	Leu	Glu	Pro	Glu	Asp	Val	Ala	Leu	95	100	105	
Tyr	Tyr	Cys	Gln	Gln	Tyr	Phe	Thr	Thr	Pro	Tyr	Thr	Phe	Gly	Gln	110	115	120	
Gly	Thr	Arg	Leu	Glu	Ile	Lys	Arg	Thr	Val	Ala	Ala	Pro	Ser	Val	125	130	135	
Phe	Ile	Phe	Pro	Pro	Ser	Asp	Glu	Gln	Leu	Lys	Ser	Gly	Thr	Ala	140	145	150	
Ser	Val	Val	Cys	Leu	Leu	Asn	Asn	Phe	Tyr	Pro	Arg	Glu	Ala	Lys	155	160	165	
Val	Gln	Trp	Lys	Val	Asp	Asn	Ala	Leu	Gln	Ser	Gly	Asn	Ser	Gln	170	175	180	

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<210> 3
 <211> 962
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> Incyte ID No: 135698CB1

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 tcacagagca ggacagcaag gacagcacct acagcctcag cagcacctg acgctgagca 660
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 aa 962

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 <212> DNA
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 <222> 2533
 <223> a or g or c or t, unknown, or other

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 gattgcgtgc gcccgcgga ggcccgggg agcggctggg atcctcagcg gcggcgggtt 180
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 gtcaccagtg acatttaatt ccaaacagga gtccttcggg ccagcaaagc tgcccaggct 840

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Ser Leu Pro Gln Leu Asn Gln Ala Ala Tyr Asp Ser Leu Val Val	125	130	135
Gly Lys Leu Ser Phe Asp Ser Ser Pro Thr Ser Ser Thr Asp Gly	140	145	150
His Ser Ser Tyr Gly Leu Asp Ser Gly Phe Cys Thr Ile Ser Arg	155	160	165
Leu Pro Arg Ser Glu Lys Pro His Asp Arg Asp Arg Asp Gly Ser	170	175	180
Phe Pro Ser Glu Pro Gly Leu Arg Arg Ser Asp Ser Leu Leu Ser	185	190	195
Phe Arg Leu Asp Leu Asp Leu Gly Pro Ser Leu Leu Ser Glu Leu	200	205	210
Leu Gly Val Met Ser Leu Pro Glu Ala Pro Ala Ala Glu Thr Pro	215	220	225
Ala Pro Ala Ala Asn Pro Pro Ala Pro Thr Ala Asn Pro Thr Gly	230	235	240
Pro Ala Ala Asn Pro Pro Ala Thr Thr Ala Asn Pro Pro Ala Pro	245	250	255
Ala Ala Asn Pro Ser Ala Pro Ala Ala Thr Pro Thr Gly Pro Ala	260	265	270
Ala Asn Pro Pro Ala Pro Ala Ala Ser Ser Thr Pro His Gly His	275	280	285
Cys Pro Asn Gly Val Thr Ala Gly Leu Gly Pro Val Ala Glu Val	290	295	300
Lys Ser Ser Pro Val Gly Gly Gly Pro Arg Gly Pro Ala Gly Pro	305	310	315
Ala Leu Gly Arg His Trp Gly Ala Gly Trp Asp Gly Gly His His	320	325	330
Tyr Pro Glu Met Asp Ala Arg Gln Glu Arg Val Glu Val Leu Pro	335	340	345
Gln Ala Arg Ala Ser Trp Glu Ser Leu Asp Glu Glu Trp Arg Ala	350	355	360
Pro Gln Ala Gly Ser Arg Thr Pro Val Pro Ser Thr Val Gln Ala	365	370	375
Asn Thr Phe Glu Phe Ala Asp Ala Glu Glu Asp Asp Glu Val Lys	380	385	390
Val			

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